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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)  
217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2008; month=1; day=11; hr=17; min=2; sec=10; ms=804; ]

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Reviewer Comments:

Please realign amino acid number so that it lines up with every fifth  
amino acid.

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Application No: 10572522

Version No: 1.0

Input Set:

Output Set:

Started: 2007-12-18 14:57:25.234

Finished: 2007-12-18 14:57:26.071

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 837 ms

Total Warnings: 0

Total Errors: 5

No. of SeqIDs Defined: 1

Actual SeqID Count: 1

Error code	Error Description
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (20)
E 323	Invalid/missing amino acid numbering SEQID (1) POS (24)
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (25)
E 323	Invalid/missing amino acid numbering SEQID (1) POS (29)
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (30)

SEQUENCE LISTING

<110> Andreas BERGMANN

<120> METHOD FOR THE DIAGNOSIS OF DISEASES BY DETERMINING APOLIPOPROTEIN C-1

<130> 2582.011

<140> 10572522

<141> 2007-12-18

<150> PCT/EP2004/09965

<151> 2004-09-07

<150> DE 103 43 815.7

<151> 2003-09-22

<160> 1

<170> PatentIn version 2.1

<210> 1

<211> 57

<212> PRT

<213> Homo sapiens

<400> 1

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Thr	Leu	Glu	Asp	Lys	Ala	Arg	Glu	Leu	Ile	Ser	Arg	Ile	Lys	Gln	Ser
							20					25			30

Glu	Leu	Ser	Ala	Lys	Met	Arg	Glu	Trp	Phe	Ser	Glu	Thr	Phe	Gln	Lys
				35				40				45			

Val	Lys	Glu	Lys	Leu	Lys	Ile	Asp	Ser
	50					55		